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OM protein - protein search, using sw model

Run on: June 27, 2003, 18:02:05 ; Search time 109 seconds
(without alignments)
35.705 Million cell updates/sec

Title: US-09-300-612-1

Perfect score: 84

Sequence: 1 LKAMDTPPLWKTE 15

Scoring table:

Gapop 10.0 , Gapext 0.5

Searched: 1184592 seqs, 259455700 residues

Total number of hits satisfying chosen parameters: 1184592

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

- 1: /cgn2_6/ptodata/1/paa/PCT_NEW_COMB.pep.*
- 2: /cgn2_6/ptodata/1/paa/US06_NEW_COMB.pep.*
- 3: /cgn2_6/ptodata/1/paa/US07_NEW_COMB.pep.*
- 4: /cgn2_6/ptodata/1/paa/US08_NEW_COMB.pep.*
- 5: /cgn2_6/ptodata/1/paa/US09_NEW_COMB.pep.*
- 6: /cgn2_6/ptodata/1/paa/US10_NEW_COMB.pep.*
- 7: /cgn2_6/ptodata/1/paa/US60_NEW_COMB.pep.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	84	100.0	15	1	PCT-US03-01044-2
2	69	82.1	12	1	PCT-US03-01044-4
3	65	77.4	11	1	PCT-US03-01044-5
4	54	64.3	10	1	PCT-US03-01044-1
5	50	59.5	9	1	PCT-US03-01044-6
6	47	56.0	531	1	PCT-US02-28538-104
7	47	56.0	531	6	US-10-236-417-104
8	46	54.8	85	6	US-10-424-599-278038
9	46	54.8	384	6	US-10-437-963-200494
10	45	53.6	41	6	US-10-424-599-173148
11	45	53.6	137	5	US-09-134-000C-5938
12	45	53.6	137	5	US-09-134-000C-5938
13	45	53.6	137	6	US-10-434-665-5938
14	44	52.4	70	6	US-10-424-599-229035
15	44	52.4	106	6	US-10-424-599-255201
16	44	52.4	124	6	US-10-424-599-151167
17	44	52.4	144	6	US-10-424-599-205702
18	44	52.4	158	6	US-10-425-114-63058
19	44	52.4	223	7	US-60-452-680-17966
20	44	52.4	372	5	US-09-724-676-66676
21	44	52.4	372	5	US-09-724-676A-66676
22	44	52.4	381	5	US-09-724-676-66678
23	44	52.4	381	5	US-09-724-676A-66678
24	44	52.4	530	6	US-10-424-599-283741
25	44	52.4	538	5	US-09-724-676-66677
26	44	52.4	538	5	US-09-724-676A-66677

ALIGNMENTS

RESULT 1

PCT-US03-01044-2
; Sequence 2, Application PC/TUS0301044

; GENERAL INFORMATION:

; APPLICANT: LIPPS, BINIE L.

; TITLE OF INVENTION: DIAGNOSIS AND TREATMENT FOR IMMUNOGLOBULIN E (IGE) IMPLICATED

; FILE REFERENCE: FWLPAT0150

; CURRENT APPLICATION NUMBER: PCT/US03/01044

; CURRENT FILING DATE: 2003-01-14

; PRIOR APPLICATION NUMBER: US 10/047,945

; PRIOR FILING DATE: 2002-01-14

; NUMBER OF SEQ ID NOS: 7

; SOFTWARE: PatentIn version 3.1

; SEQ ID NO 2

; LENGTH: 15

; TYPE: PRT

; ORGANISM: ARTIFICIAL

; FEATURE:

; OTHER INFORMATION: SYNTHETIC FRAGMENT OF ISOLATE FROM OPOSSUM SERUM. SEE US 5,57

; OTHER INFORMATION: 97.

PCT-US03-01044-2

Query Match 100.0%; Score 84; DB 1; Length 15;
Best Local Similarity 100.0%; Pred. No. 2.1e-05;
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LKAMDTPPLWKTE 15

Db 1 LKAMDTPPLWKTE 15

RESULT 2

PCT-US03-01044-4
; Sequence 4, Application PC/TUS0301044

; GENERAL INFORMATION:

; APPLICANT: LIPPS, BINIE L.

; TITLE OF INVENTION: DIAGNOSIS AND TREATMENT FOR IMMUNOGLOBULIN E (IGE) IMPLICATED

; FILE REFERENCE: FWLPAT0150

; CURRENT APPLICATION NUMBER: PCT/US03/01044

; CURRENT FILING DATE: 2003-01-14

; PRIOR APPLICATION NUMBER: US 10/047,945

; PRIOR FILING DATE: 2002-01-14

; NUMBER OF SEQ ID NOS: 7

; SOFTWARE: PatentIn version 3.1

; SEQ ID NO 4

; LENGTH: 12

; TYPE: PRT

Sequence 66675, A
Sequence 66675, A
Sequence 238, App
Sequence 9380, App
Sequence 7, Appli
Sequence 168841,
Sequence 114167,
Sequence 197262,
Sequence 197265,
Sequence 51474, A
Sequence 73022, A
Sequence 75158, A
Sequence 75894, A
Sequence 566, App
Sequence 144200,
Sequence 128480,
Sequence 264272,
Sequence 14216, A
Sequence 14216, A

US-09-724-676-66675
PCT-US02-19669A-228
US-10-325-899-9380
PCT-US03-01044-7
US-10-437-963-168841
US-10-437-963-114167
US-10-437-963-197262
US-10-437-963-197265
US-10-282-122A-51474
US-10-282-122A-73022
US-10-282-122A-75158
US-10-282-122A-75894
US-10-361-942-566
US-10-437-963-144200
US-10-437-963-128480
US-10-424-599-264272
PCT-US02-32727-14216
US-09-978-825-14216

52.4 679 5
52.4 679 5
52.4 1336 1
52.4 1336 6
52.4 1336 8
51.2 129 6
51.2 156 6
51.2 258 6
51.2 278 6
51.2 286 6
51.2 341 6
51.2 341 6
51.2 341 6
51.2 687 6
51.2 840 6
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50.0 92 6
50.0 108 1
50.0 108 5

; ORGANISM: ARTIFICIAL
; FEATURE:
; OTHER INFORMATION: SYNTHETIC. CORRESPONDS TO FRAGMENT 1-12 OF 2 ABOVE.
PCT-US03-01044-4

Query Match 82.1%; Score 69; DB 1; Length 12;
Best Local Similarity 100.0%; Pred. No. 0.0025;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LKAMDTPPLWI 12
Db 1 LKAMDTPPLWI 12

RESULT 3
PCT-US03-01044-5
; Sequence 5, Application PC/TUS0301044
; GENERAL INFORMATION:
; APPLICANT: LIPPS, BINIE L.
; TITLE OF INVENTION: DIAGNOSIS AND TREATMENT FOR IMMUNOGLOBULIN E (IGE) IMPLICATED DIS
; FILE REFERENCE: FWLPAT015WO
; CURRENT APPLICATION NUMBER: PCT/US03/01044
; CURRENT FILING DATE: 2003-01-14
; PRIOR APPLICATION NUMBER: US 10/047,945
; PRIOR FILING DATE: 2002-01-14
; NUMBER OF SEQ ID NOS: 7
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5
; LENGTH: 11
; TYPE: PRT
; ORGANISM: ARTIFICIAL
; FEATURE:
; OTHER INFORMATION: SYNTHETIC. CORRESPONDS TO FRAGMENT 1-11 OF 2 ABOVE.
PCT-US03-01044-5

Query Match 77.4%; Score 65; DB 1; Length 11;
Best Local Similarity 100.0%; Pred. No. 0.009;
Matches 11; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LKAMDTPPLW 11
Db 1 LKAMDTPPLW 11

RESULT 4
PCT-US03-01044-1
; Sequence 1, Application PC/TUS0301044
; GENERAL INFORMATION:
; APPLICANT: LIPPS, BINIE L.
; TITLE OF INVENTION: DIAGNOSIS AND TREATMENT FOR IMMUNOGLOBULIN E (IGE) IMPLICATED DIS
; FILE REFERENCE: FWLPAT015WO
; CURRENT APPLICATION NUMBER: PCT/US03/01044
; CURRENT FILING DATE: 2003-01-14
; PRIOR APPLICATION NUMBER: US 10/047,945
; PRIOR FILING DATE: 2002-01-14
; NUMBER OF SEQ ID NOS: 7
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 1
; LENGTH: 10
; TYPE: PRT
; ORGANISM: ARTIFICIAL
; FEATURE:
; OTHER INFORMATION: SYNTHETIC FRAGMENT OF ISOLATE FROM OPOSSUM SERUM. SEE US 5,576,2
; OTHER INFORMATION: 87.
PCT-US03-01044-1

Query Match 64.3%; Score 54; DB 1; Length 10;
Best Local Similarity 100.0%; Pred. No. 0.33;
Matches 10; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LKAMDTPPL 10

Db 1 LKAMDTPPL 10

RESULT 5
PCT-US03-01044-6
; Sequence 6, Application PC/TUS0301044
; GENERAL INFORMATION:
; APPLICANT: LIPPS, BINIE L.
; TITLE OF INVENTION: DIAGNOSIS AND TREATMENT FOR IMMUNOGLOBULIN E (IGE) IMPLICATED
; FILE REFERENCE: FWLPAT015WO
; CURRENT APPLICATION NUMBER: PCT/US03/01044
; CURRENT FILING DATE: 2003-01-14
; PRIOR APPLICATION NUMBER: US 10/047,945
; PRIOR FILING DATE: 2002-01-14
; NUMBER OF SEQ ID NOS: 7
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 6
; LENGTH: 9
; TYPE: PRT
; ORGANISM: ARTIFICIAL
; FEATURE:
; OTHER INFORMATION: SYNTHETIC. CORRESPONDS TO FRAGMENT 1-9 OF 2 ABOVE.
PCT-US03-01044-6

Query Match 59.5%; Score 50; DB 1; Length 9;
Best Local Similarity 100.0%; Pred. No. 1.1e+06;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LKAMDTPPP 9
Db 1 LKAMDTPPP 9

RESULT 6
PCT-US02-28538-104
; Sequence 104, Application PC/TUS0228538
; GENERAL INFORMATION:
; APPLICANT: Curagen Corp. et al.
; TITLE OF INVENTION: NOVEL PROTEINS AND NUCLEIC ACIDS ENCODING SAME
; FILE REFERENCE: 21402-442C
; CURRENT APPLICATION NUMBER: PCT/US02/28538
; CURRENT FILING DATE: 2002-12-04
; PRIOR APPLICATION NUMBER: US60/318,120
; PRIOR FILING DATE: 2001-09-01
; PRIOR APPLICATION NUMBER: US60/318,430
; PRIOR FILING DATE: 2001-09-10
; PRIOR APPLICATION NUMBER: US60/322,781
; PRIOR FILING DATE: 2001-09-17
; PRIOR APPLICATION NUMBER: US60/318,184
; PRIOR FILING DATE: 2001-09-07
; PRIOR APPLICATION NUMBER: US60/361,663
; PRIOR FILING DATE: 2002-03-05
; PRIOR APPLICATION NUMBER: US60/396,412
; PRIOR FILING DATE: 2002-07-17
; PRIOR APPLICATION NUMBER: US60/322,636
; PRIOR FILING DATE: 2001-09-17
; PRIOR APPLICATION NUMBER: US60/322,817
; PRIOR FILING DATE: 2001-09-17
; PRIOR APPLICATION NUMBER: US60/322,816
; PRIOR FILING DATE: 2001-09-17
; PRIOR APPLICATION NUMBER: US60/323,519
; PRIOR FILING DATE: 2001-09-19
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 341
; SOFTWARE: Custom
; SEQ ID NO 104
; LENGTH: 531
; TYPE: PRT
; ORGANISM: Homo sapiens
PCT-US02-28538-104

Query Match 56.0%; Score 47; DB 1; Length 531;
Best Local Similarity 66.7%; Pred. No. 1.1e+02;
Matches 8; Conservative 1; Mismatches 3; Indels 0; Gaps 0;

QY 4 MDPTPLWIKTE 15
||:|||||
Db 70 MDTSPPLWLTLE 81

RESULT 7

US-10-236-417-104
; Sequence 104, Application US/10236417
; GENERAL INFORMATION:
; APPLICANT: Agee et al.
; TITLE OF INVENTION: NOVEL PROTEINS AND NUCLEIC ACIDS ENCODING SAME
; FILE REFERENCE: 21402-442C
; CURRENT APPLICATION NUMBER: US/10/236,417
; CURRENT FILING DATE: 2003-01-06
; PRIOR APPLICATION NUMBER: US60/318,120
; PRIOR FILING DATE: 2001-09-01
; PRIOR APPLICATION NUMBER: US60/318,430
; PRIOR FILING DATE: 2001-09-10
; PRIOR APPLICATION NUMBER: US60/322,781
; PRIOR FILING DATE: 2001-09-17
; PRIOR APPLICATION NUMBER: US60/318,184
; PRIOR FILING DATE: 2001-09-07
; PRIOR APPLICATION NUMBER: US60/361,663
; PRIOR FILING DATE: 2002-03-05
; PRIOR APPLICATION NUMBER: US60/396,412
; PRIOR FILING DATE: 2002-07-17
; PRIOR APPLICATION NUMBER: US60/322,636
; PRIOR FILING DATE: 2001-09-17
; PRIOR APPLICATION NUMBER: US60/322,817
; PRIOR FILING DATE: 2001-09-17
; PRIOR APPLICATION NUMBER: US60/322,816
; PRIOR FILING DATE: 2001-09-17
; PRIOR APPLICATION NUMBER: US60/323,519
; PRIOR FILING DATE: 2001-09-19
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 341
; SOFTWARE: Custom
; SEQ ID NO 104
; LENGTH: 531
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-236-417-104

Query Match 56.0%; Score 47; DB 6; Length 531;
Best Local Similarity 66.7%; Pred. No. 1.1e+02;
Matches 8; Conservative 1; Mismatches 3; Indels 0; Gaps 0;

QY 4 MDPTPLWIKTE 15
||:|||||
Db 70 MDTSPPLWLTLE 81

RESULT 8

US-10-424-599-278038
; Sequence 278038, Application US/10424599
; GENERAL INFORMATION:
; APPLICANT: La Rosa Thomas J
; APPLICANT: Kovalic David K
; APPLICANT: Zhou Yihua
; APPLICANT: Cao Yongwei
; TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53223)B
; CURRENT APPLICATION NUMBER: US/10/424,599
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 285684
; SEQ ID NO 278038
; LENGTH: 85
; TYPE: PRT

; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT3847_93091C.1.pep
US-10-424-599-278038

Query Match 54.8%; Score 46; DB 6; Length 85;
Best Local Similarity 77.8%; Pred. No. 31;
Matches 7; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 7 TPPLWIKTE 15
|||||
Db 45 TPPLWVKIE 53

RESULT 9

US-10-437-963-200494
; Sequence 200494, Application US/10437963
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; APPLICANT: Wu, Wei
; APPLICANT: Boukharov, Andrey A.
; APPLICANT: Barbazuk, Brad
; APPLICANT: Li, Ping
; TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53221)B
; CURRENT APPLICATION NUMBER: US/10/437,963
; CURRENT FILING DATE: 2003-05-14
; NUMBER OF SEQ ID NOS: 204966
; SEQ ID NO 200494
; LENGTH: 384
; TYPE: PRT
; ORGANISM: Oryza sativa
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT4530_95958C.1.pep
US-10-437-963-200494

Query Match 54.8%; Score 46; DB 6; Length 384;
Best Local Similarity 58.3%; Pred. No. 1.1e+02;
Matches 7; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

QY 3 AMDPTPLWIKT 14
|:|||||
Db 91 AAAPAPPLWLRT 102

RESULT 10

US-10-424-599-173148
; Sequence 173148, Application US/10424599
; GENERAL INFORMATION:
; APPLICANT: La Rosa Thomas J
; APPLICANT: Kovalic David K
; APPLICANT: Zhou Yihua
; APPLICANT: Cao Yongwei
; TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53223)B
; CURRENT APPLICATION NUMBER: US/10/424,599
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 285684
; SEQ ID NO 173148
; LENGTH: 41
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT3847_127369C.1.pep
US-10-424-599-173148

Query Match 53.6%; Score 45; DB 6; Length 41;
Best Local Similarity 63.6%; Pred. No. 23;

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Matches 7; Conservative 1; Mismatches 3; Indels 0; Gaps 0;

Qy 5 DPTPLWIKTE 15
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Db 14 DPPLWAOPE 24

RESULT 11
US-09-134-000C-5938
; Sequence 5938, Application US/09134000C
; GENERAL INFORMATION:
; APPLICANT: Lynn Doucette-Stamm et al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
; FILE OF INVENTION: ENTEROCOCCUS FAECALIS FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: 032796-032
; CURRENT APPLICATION NUMBER: US/09/134,000C
; CURRENT FILING DATE: 1998-08-13
; PRIOR APPLICATION NUMBER: US 60/055,778
; PRIOR FILING DATE: 1997-08-15
; NUMBER OF SEQ ID NOS: 6812
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5938
; LENGTH: 137
; TYPE: PRT
; ORGANISM: Enterococcus faecalis
US-09-134-000C-5938

Query Match 53.6%; Score 45; DB 5; Length 137;
Best Local Similarity 75.0%; Pred. No. 65;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

Qy 5 DPTPLW 12
   :|||||
Db 85 NPTPLW 92

RESULT 12
US-09-134-000C-5938
; Sequence 5938, Application US/09134000C
; GENERAL INFORMATION:
; APPLICANT: Lynn Doucette-Stamm et al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
; FILE OF INVENTION: ENTEROCOCCUS FAECALIS FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: 032796-032
; CURRENT APPLICATION NUMBER: US/09/134,000C
; CURRENT FILING DATE: 1998-08-13
; PRIOR APPLICATION NUMBER: US 60/055,778
; PRIOR FILING DATE: 1997-08-15
; NUMBER OF SEQ ID NOS: 6812
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5938
; LENGTH: 137
; TYPE: PRT
; ORGANISM: Enterococcus faecalis
US-09-134-000C-5938

Query Match 53.6%; Score 45; DB 5; Length 137;
Best Local Similarity 75.0%; Pred. No. 65;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

Qy 5 DPTPLW 12
   :|||||
Db 85 NPTPLW 92

RESULT 13
US-09-134-000C-5938
; Sequence 5938, Application US/09134000C
; GENERAL INFORMATION:
; APPLICANT: Lynn Doucette-Stamm et al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
; FILE OF INVENTION: ENTEROCOCCUS FAECALIS FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: PATH03-09
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; CURRENT APPLICATION NUMBER: US/10/434,665
; CURRENT FILING DATE: 2003-05-14
; PRIOR APPLICATION NUMBER: US 09/134,000
; PRIOR FILING DATE: 1998-08-13
; PRIOR APPLICATION NUMBER: US 60/055,778
; PRIOR FILING DATE: 1997-08-15
; NUMBER OF SEQ ID NOS: 6812
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5938
; LENGTH: 137
; TYPE: PRT
; ORGANISM: Enterococcus faecalis
US-10-434-665-5938

Query Match 53.6%; Score 45; DB 6; Length 137;
Best Local Similarity 75.0%; Pred. No. 65;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

Qy 5 DPTPLW 12
   :|||||
Db 85 NPTPLW 92

RESULT 14
US-10-424-599-229035
; Sequence 229035, Application US/10424599
; GENERAL INFORMATION:
; APPLICANT: La Rosa Thomas J
; APPLICANT: Kovalic David K
; APPLICANT: Zhou Yihua
; APPLICANT: Cao Yongwei
; TITLE OF INVENTION: SOY NUCLEIC ACID MOLECULES AND OTHER MOLECULES ASSOCIATED WITH
; FILE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(53223)B
; CURRENT APPLICATION NUMBER: US/10/424,599
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 285684
; SEQ ID NO 229035
; LENGTH: 70
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT3847_48845C.1.pep
US-10-424-599-229035

Query Match 52.4%; Score 44; DB 6; Length 70;
Best Local Similarity 60.0%; Pred. No. 51;
Matches 9; Conservative 0; Mismatches 6; Indels 0; Gaps 0;

Qy 1 LKAMDPTPLWIKTE 15
   |||||
Db 55 LKARDQOKSEWIKTE 69

RESULT 15
US-10-424-599-255201
; Sequence 255201, Application US/10424599
; GENERAL INFORMATION:
; APPLICANT: La Rosa Thomas J
; APPLICANT: Kovalic David K
; APPLICANT: Zhou Yihua
; APPLICANT: Cao Yongwei
; TITLE OF INVENTION: SOY NUCLEIC ACID MOLECULES AND OTHER MOLECULES ASSOCIATED WITH
; FILE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(53223)B
; CURRENT APPLICATION NUMBER: US/10/424,599
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 285684
; SEQ ID NO 255201
; LENGTH: 106
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
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; OTHER INFORMATION: Clone ID: PAT_MRT3847_72469C.1.pap
US-10-424-599-255201

Query Match 52.4%; Score 44; DB 6; Length 106;
Best Local Similarity 85.7%; Pred. No. 73;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

OY 6 PTPPLWI 12
| | | | |
Db 92 PTPPLWV 98

Search completed: June 27, 2003, 18:10:15
Job time : 111 secs

c.